REMARKS/ARGUMENTS

Status of the Claims

Upon entry of the present amendment, claims 4-8 and 15-23 are pending. Claims 4-6 are amended. New claims 15-23 are added.

Claims 4-6 are amended to set forth specific concentrations of HGF and FGF-2. Support is found, for example, on page 16, line 24 through page 17, line 1.

New claims 15 and 16 find support, for example, on page 16, line 24 through page 17, line 1.

New claim 17 finds support, for example, on page 29, lines 15-16. New claim 18 finds support, for example, on page 37, line 16 through page 38,

line 6.

New claims 19-21 find support, for example, on page 29, lines 3-22 and throughout the specification. It is clear that the culturing conditions of the present invention were carried out under atmospheric oxygen levels.

New claim 22, finds support, for example, on page 8, lines 17-20.

New claim 23 finds support, for example, on page 8, lines 21-22.

The present amendments are necessary to place the claims in form for allowance or to reduce issues for appeal. No new matter is added by the present amendments, and the Examiner is respectfully requested to enter them.

Claim Objections

The Examiner objected to claims 4-6 for reciting non-elected subject matter. This objection is obviated by amendment of claims 4-6 to cancel recitation of non-elected subject matter.

Rejections under 35 U.S.C. § 112, second paragraph

The Examiner has rejected claims 6-8 under 35 U.S.C. § 112, second paragraph, as allegedly unclear. Applicants do not agree with the Examiner. However, in the interest of

furthering prosecution, Applicants have amended claim 6 to set forth the step of culturing cells under conditions that allow their differentiation into a population of cells containing neurons and glia.

Rejection under 35 U.S.C. § 102(e)

The Examiner has maintained the rejection of claims 4-6 and 8 under 35 U.S.C. § 102(e) as allegedly anticipated by U.S. Patent No. 6,589,728 ("Csete"). Applicants do not agree with the Examiner. However, in the interest of furthering prosecution, Applicants have amended claims 4-6 to set forth particular concentrations of HGF and FGF-2. Csete does not teach or suggest particularly using HGF and FGF-2 to culture, proliferate or differentiate neural stem cells, much less teach or suggest a particular concentration range for either growth factor. Furthermore, the invention in Csete relies on subatmospheric oxygen levels in culture (i.e., less than 12% in the claims). This is reflected in the abstract and claims of Csete.

The Examiner is respectfully reminded that the present invention is a selection invention particularly directed to the culture, proliferation and differentiation of <u>neural</u> stem cells. The passages in columns 7 and 15 of Csete identified by the Examiner are not entirely consistent with each other. The passage at column 7, lines 42-62 is concerned with <u>stem cells generally</u> and not neural stem cells in particular. The passage at column 7 does not teach or suggest particularly combining HGF with FGF-2 (a.k.a., bFGF) to culture, proliferate or differentiate any kind of stem cell, much less particularly neural stem cells.

The passage at column 15, lines 51-65 of Csete is expressly directed to isolating and culturing neural stem cells. Csete discloses that neuroepithelial stem cells can be cultured and proliferated in FGF-2 (bFGF), but does not disclose or suggest combining FGF-2 with HGF. With respect to differentiation of neural stem cells into neurons and glia, Csete affirmatively states that the FGF-2 (bFGF) is removed and replaced with media lacking FGF-2. Therefore, in the passage at column 15, lines 63-65, Csete expressly teaches that differentiation of neural stem cells is performed in the absence of FGF-2 and does not teach or suggest adding HGF for the purpose of culturing, proliferating or differentiating neural stem cells. The Examiner can not

give greater weight to the general disclosure of Csete in column 7 while ignoring the disclosure particular to neural stem cells at column 15.

Because Csete does not teach or suggest each and every element of the claimed methods, Csete does not anticipate the present invention. Accordingly, the Examiner is respectfully requested to withdraw this rejection.

Rejection under 35 U.S.C. § 103(a)

The Examiner has rejected claim 7 under 35 U.S.C. § 103(a) as allegedly obvious over Csete in view of U.S. Patent No. 5,753,505 ("Luskin"). To the extent that the present rejection applies to the present claims, Applicants respectfully traverse.

Applicants maintain the position that the combined disclosures of Csete and Luskin do not teach or suggest all of the steps and elements of the claimed methods. The Examiner is respectfully reminded that the present methods are a selection invention particularly directed to the culture, proliferation and differentiation of neural stem cells by culturing them in a growth medium comprising the particularly selected combination of HGF and FGF-2. As discussed previously, in the passage in column 7 of Csete, no particular combination of growth factors is called out and no particular stem cell type is called out. As a matter of enablement of Csete as a reference, the skilled person is still left to determine which out of the myriad of permutations of growth factors and stem cell types to match up for the culture, proliferation and/or differentiation conditions particular to neural stem cells. Where Csete does expressly discuss neural stem cells in the passage in column 15, Csete teaches away from the present methods by teaching that FGF-2 is removed prior to differentiation neural stem cells. Csete makes absolutely no mention of HGF in the growth medium of neural stem cells.

With respect to the amended claims, Csete does not teach or suggest the particular concentrations of growth factors. Csete certainly teaches against the use of atmospheric concentrations of oxygen in culturing, proliferating, or differentiating any kind of stem cell, including neural stem cells. See, e.g., abstract, summary and claims of Csete. As stated previously, Luskin does not cure the deficiencies of Csete. Luskin discloses adding nerve

growth factor (NGF) or brain-derived neurotrophic factor (BDNF) to the growth medium of neural stem cells, but does not teach or suggest in any way that <u>hepatocyte growth factor</u> would find use in culturing, proliferating or differentiation <u>neural stem cells</u>. Based on the disclosures of Csete and Luskin, the skilled person would have no reason to expect that a liver cell growth factor should promote differentiation of nerve stem cells, for example, into neurons.

In any case, Applicants have rebutted any alleged prima facie case of obviousness by demonstrating an unexpected synergistic effect of HGF and FGF-2 in promoting the growth and differentiation of neural stem cells. This is shown in columns 1-3 of Table 1 on page 35 of the present application. The Examiner alleges that the synergistic effects of HGF and FGF2 are not unexpected because HGF and FGF-2 are structurally and functionally distinct growth factors.

See, pages 4-5 of the present Office Action. However, FGF-2 and EGF also are structurally and functionally distinct growth factors, and their combined effects were less than additive. See, columns 2 (FGF-2 only), 4 (EGF only) and 6 (FGF-2 and EGF) of Table 1. There is no a priori reason for the skilled person to expect that the combination of HGF and FGF-2 should be synergistic and the combination of FGF-2 and EGF be less than additive in promoting the proliferation of neural stem cells.

Also, the differentiated neural cell populations are different in the presence or absence of HGF. In the presence of HGF, a majority of the differentiated cells are neurons. In the presence of EGF and FGF-2, but in the absence of HGF, only about 30% of the differentiated cells are neurons. See, e.g., page 37, lines 16-24 of the present application.

In view of the foregoing, the combined disclosures of Csete and Luskin do not render the present methods obvious. Accordingly, the Examiner is respectfully requested to withdraw the present rejection.

Amino acid sequences and BLAST alignments of human HGF, FGF-2 and EGF are attached as Exhibit A.

PATENT

Appl. No. 10/536,563 Amdt. dated December 21, 2007 Amendment under 37 CFR 1.116 Expedited Procedure Examining Group 1633

CONCLUSION

In view of the foregoing, Applicants believe all claims now pending in this

Application are in condition for allowance and an action to that end is respectfully requested.

If the Examiner believes a telephone conference would expedite prosecution of

this application, please telephone the undersigned at 415-576-0200.

Respectfully submitted,

Jumily Wallett

Jennifer L. Wahlsten Reg. No. 46,226

TOWNSEND and TOWNSEND and CREW LLP Two Embarcadero Center, Eighth Floor San Francisco, California 94111-3834 Tel: 415-576-0200 Fax: 415-576-0300 Attachments ILW:ilw

61201308 v1

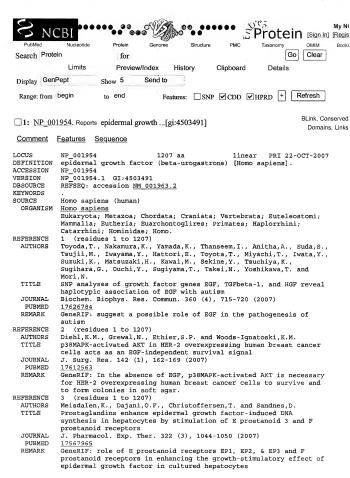


Exhibit A

```
REFERENCE
            4 (residues 1 to 1207)
  AUTHORS
            Su, X., Kong, C. and Stahl, P.D.
  TITLE
          GAPex-5 mediates ubiquitination, trafficking, and degradation of
            epidermal growth factor receptor
  JOURNAL
            J. Biol. Chem. 282 (29), 21278-21284 (2007)
   PUBMED
            17545148
  REMARK
            GeneRIF: EGF-stimulated receptor ubiquitination and trafficking are
            mediated via GAPex-5: GAPex-5-mediated EGFR ubiquitination is
            independent of Rab5 activation
REFERENCE 5 (residues 1 to 1207)
  AUTHORS
            Durer, U., Hartig, R., Bang, S., Thim, L. and Hoffmann, W.
  TITLE
            TFF3 and EGF induce different migration patterns of intestinal
            epithelial cells in vitro and trigger increased internalization of
            E-cadherin
  TOTTRNAT.
            Cell. Physiol. Biochem. 20 (5), 329-346 (2007)
   PUBMED
            17762162
  REMARK
            GeneRIF: trefoil factor 3, in contrast to EGF, enhanced a
            collective cell migration ensuring a precise coverage of the
            re-populated area avoiding gaps
REFERENCE
            6 (sites)
  AUTHORS
           Skidgel, R.A., McGwire, G.B. and Li, X.Y.
  TITLE
           Membrane anchoring and release of carboxypeptidase M: implications
            for extracellular hydrolysis of peptide hormones
  JOURNAL
           Immunopharmacology 32 (1-3), 48-52 (1996)
   PUBMED
            8796265
REFERENCE
           7
               (sites)
  AUTHORS
           McGwire, G.B. and Skidgel, R.A.
  TITLE
            Extracellular conversion of epidermal growth factor (EGF) to
           des-Arq53-EGF by carboxypeptidase M
  JOURNAL J. Biol. Chem. 270 (29), 17154-17158 (1995)
   DITEMED
          7615511
REFERENCE
            8 (residues 1 to 1207)
  AUTHORS Ishibashi, T., Bottaro, D.P., Chan, A., Miki, T. and Aaronson, S.A.
           Expression cloning of a human dual-specificity phosphatase
  TITLE
  JOURNAL Proc. Natl. Acad. Sci. U.S.A. 89 (24), 12170-12174 (1992)
   PUBMED 1281549
REFERENCE
           9 (residues 1 to 1207)
  AUTHORS
           Gout, I., Dhand, R., Panayotou, G., Fry, M.J., Hiles, I., Otsu, M. and
            Waterfield, M.D.
  TITLE
            Expression and characterization of the p85 subunit of the
            phosphatidylinositol 3-kinase complex and a related p85 beta
           protein by using the baculovirus expression system
  JOURNAL
           Biochem. J. 288 (PT 2), 395-405 (1992)
   PURMED
            1334406
REFERENCE
            10 (residues 1 to 1207)
  AUTHORS
           Hommel, U., Harvey, T.S., Driscoll, P.C. and Campbell, I.D.
  TITLE
           Human epidermal growth factor. High resolution solution structure
           and comparison with human transforming growth factor alpha
  JOURNAL
           J. Mol. Biol. 227 (1), 271-282 (1992)
   PUBMED
          1522591
REFERENCE
           11 (residues 1 to 1207)
  AUTHORS
            Lei, Z.M. and Rao, C.V.
  TITLE
            Expression of epidermal growth factor (EGF) receptor and its
            ligands, EGF and transforming growth factor-alpha, in human
            fallopian tubes
  JOURNAL
          Endocrinology 131 (2), 947-957 (1992)
   PUBMED 1639032
REFERENCE 12 (residues 1 to 1207)
  AUTHORS Gregory, H. and Preston, B.M.
  TITLE
           The primary structure of human urogastrone
```

JOURNAL PUBMED Int. J. Pept. Protein Res. 9 (2), 107-118 (1977)
300079

COMMENT REVIEWS

REVIEWED REFSEQ: This record has been curated by NCBI staff. The reference sequence was derived from X04571.1 and AF023155.1.

Summary: Bpidermal growth factor has a profound effect on the differentiation of specific cells in vivo and is a potent mitogenic factor for a variety of cultured cells of both ectodermal and mesodermal origin. The EGF precursor is believed to exist as a membrane-bound molecule which is proteolytically cleaved to generate the 53-amino acid peptide hormone that stimulates cells to divide.

Publication Note: This RefSeq record includes a subset of the publications that are available for this gene. Please see the Entrez Gene record to access additional publications.

FEATURES

Location/Qualifiers
1..1207

source

/organism="Homo sapiens" /db_xref="taxon:9606"

/chromosome="4" /map="4q25"

Protein

1..1207
/product="epidermal growth factor (beta-urogastrone)"

/note="urogastrone"
1..22

sig peptide

/calculated mol wt=2392

Region

151..189

/region_name="LY"

/note="Low-density lipoprotein-receptor YWTD domain; Type 'B' repeats in low-density lipoprotein (LDL) receptor that plays a central role in mammalian cholesterol metabolism.

Also present in a variety of molecules similar to

gp300/megalin; smart00135"

/db_xref="CDD:47474"

Region

<352..393 /region name="vWA Matrilin"

/note="TWNA_Matrilin: In cartilaginous plate, extracellular matrix molecules mediate cell-matrix and matrix-matrix interactions thereby providing tissue integrity. Some members of the matrilin family are expressed specifically

in developing cartilage rudiments; cd01475"

/db xref="CDD:29248"

/db_xrer="CDD:<u>29248</u>" 505..546

Region 505

/region_name="LY"

/note="Low-density lipoprotein-receptor YWTD domain; Type 'B' repeats in low-density lipoprotein (LDL) receptor that plays a central role in mammalian cholesterol metabolism.

Also present in a variety of molecules similar to

gp300/megalin; smart00135"

/db_xref="CDD:47474"

Region 547..587

/region name="LY"

/note="Low-density lipoprotein-receptor YWTD domain; Type 'B' repeats in low-density lipoprotein (LDL) receptor that plays a central role in mammalian cholesterol metabolism.

Also present in a variety of molecules similar to

qp300/megalin; smart00135"

/db xref="CDD:47474"

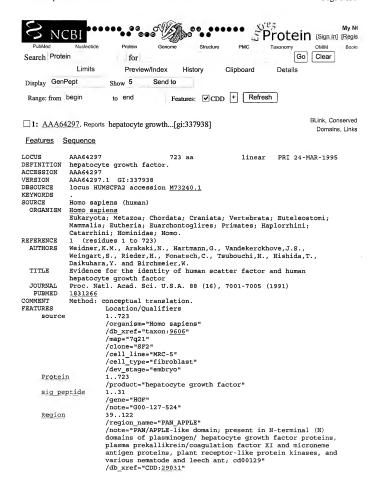
Region 590..633

```
/region name="LY"
                /note="Low-density lipoprotein-receptor YWTD domain; Type
                'B' repeats in low-density lipoprotein (LDL) receptor that
                plays a central role in mammalian cholesterol metabolism.
                Also present in a variety of molecules similar to
                qp300/megalin; smart00135"
                /db xref="CDD:47474"
Region
                635..676
                /region name="LY"
                /note="Low-density lipoprotein-receptor YWTD domain; Type
                'B' repeats in low-density lipoprotein (LDL) receptor that
                plays a central role in mammalian cholesterol metabolism.
                Also present in a variety of molecules similar to
                qp300/megalin; smart00135"
                /db_xref="CDD:47474"
                677..718
Region
                /region name="LY"
                /note="Low-density lipoprotein-receptor YWTD domain; Type
                'B' repeats in low-density lipoprotein (LDL) receptor that
                plays a central role in mammalian cholesterol metabolism.
                Also present in a variety of molecules similar to
                gp300/megalin; smart00135"
                /db xref="CDD:47474"
Region
                <740..779
                /region name="vWA Matrilin"
                /note="VWA Matrilin: In cartilaginous plate, extracellular
                matrix molecules mediate cell-matrix and matrix-matrix
                interactions thereby providing tissue integrity. Some
                members of the matrilin family are expressed specifically
                in developing cartilage rudiments; cd01475"
                /db xref="CDD:29248"
                870..910
Region
                /region name="EGF CA"
                /note="Calcium-binding EGF-like domain; smart00179"
                /db xref="CDD:47510"
Region
                912..>940
                /region name="EGF CA"
                /note="Calcium binding EGF domain; pfam07645"
                /db xref="CDD:71088"
mat peptide
                971..1023
                /product="epidermal growth factor"
                /calculated mol wt=6222
Site
                1022
                /site type="modified"
                /experiment="experimental evidence, no additional details
                recorded"
                /note="proteolytic cleavage site"
                /citation=[6]
                /citation=[7]
                /db xref="HPRD:00273"
CDS
                1..1207
                /gene="EGF"
                /coded by="NM 001963.2:443..4066"
                /GO component="extracellular region [PMID 9712850];
                integral to membrane; nucleus; plasma membrane [PMID
                3491360] "
                /GO function="calcium ion binding; epidermal growth factor
                receptor activating ligand activity [PMID 9712850]; growth
                factor activity [PMID 7736574]; protein binding [PMID
                2790960] "
```

```
/GO_process="activation of MAPK activity [PMID 9482941];
                     chromosome organization and biogenesis (sensu Eukaryota)
                     [PMID 10436156]; DNA replication [PMID 9482941]; epidermal
                     growth factor receptor signaling pathway; positive
                     regulation of cell proliferation [PMID 7736574]"
                     /db xref="CCDS:CCDS3689.1"
                     /db xref="GeneID:1950"
                     /db xref="HGNC:3229"
                     /db xref="HPRD:00578"
                     /db xref="MIM:131530"
ORIGIN
        1 mlltliillp vyskfsfysl sapphwscpe gtlagngnst cygpapflif shgnsifrid
       61 tegtnyeqlv vdagvsvimd fhynekriyw vdlerqllqr vflngsrqer vcnieknvsg
      121 mainwineev iwsnggegii tytdmkgnns hillsalkyp anvaydpyer fifwsseyag
      181 slyradldgy gykalletse kitaysldyl dkrlfwigyn regsnslics cdydggsyhi
      241 skhptqhnlf amslfqdrif ystwkmktiw iankhtqkdm vrinlhssfv plqelkvvhp
      301 lagpkaeddt wepegkicki rkgncsstyc ggdlgshicm caegyalsrd rkycedynec
      361 afwnhqctlq ckntpqsyyc tcpvqfvllp dqkrchqlvs cprnvsecsh dcvltseqpl
      421 cfcpegsvle rdgktcsgcs spdnggcsql cvplspvswe cdcfpgydlq ldekscaasg
      481 papfilfans gdirhmhfdg tdygtllsgg mgmyyaldhd pyenkiyfah talkwieran
      541 mdgsgrerli eegvdvpegl avdwigrrfy wtdrgkslig rsdlngkrsk iitkenisqp
      601 rgiavhpmak rlfwtdtgin priessslqg lgrlviassd liwpsgitid fltdklywcd
      661 akgsvieman ldgskrrrlt qndvghpfav avfedyvwfs dwampsvirv nkrtgkdrvr
      721 lggsmlkpss lvvvhplakp gadpclygng gcehickkrl gtawcscreg fmkasdgktc
      781 laldqhqlla ggevdlknqv tpldilsktr vsednitesq hmlvaeimvs dqddcapvqc
      841 smyarciseg edatcqclkg fagdgklcsd idecemgvpv cppasskcin teggyvcrcs
      901 egygddihc ldidecglgv hscgenasct nteggytcmc agrlsepgli cpdstppphl
      961 reddhhysvr nsdsecplsh dgyclhdgvc myiealdkya cncvvgyige rcqyrdlkww
     1021 elrhaqhqqq qkvivvavcv vvlvmlllls lwqahyyrtq kllsknpknp yeessrdvrs
     1081 rrpadtedqm sscpqpwfvv ikehqdlkng qqpvaqedqq aadqsmqpts wrqepqlcqm
     1141 gteggcwipy ssdkgscpgy mersfhmpsy gtgtleggve kphsllsanp lwggraldpp
     1201 hgmeltg
11
```

Disclaimer | Write to the Help Desk NCBI | NLM | NIH

Aug 28 2007 16:53:42



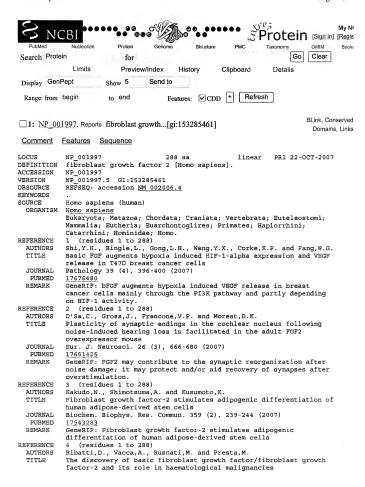
```
Site
                order (60,62,73)
                /site type="other"
                /note="putative binding site"
                /db xref="CDD:29031"
Region
                126..202
                /region name="KR"
                /note="Kringle domain; Kringle domains are believed to
                play a role in binding mediators, such as peptides, other
                proteins, membranes, or phospholipids; cd00108"
                /db xref="CDD:29008"
Site
                137
                /site type="other"
                /note="putative domain interaction site"
                /db xref="CDD:29008"
Site
                order (152, 162, 183, 185, 193)
                /site_type="other"
                /note="ligand binding site"
                /db xref="CDD:29008"
Region
                203..284
                /region name="KR"
                /note="Kringle domain; Kringle domains are believed to
                play a role in binding mediators, such as peptides, other
                proteins, membranes, or phospholipids; cd00108"
                /db xref="CDD:29008"
Site
                215
                /site_type="other"
                /note="putative domain interaction site"
                /db xref="CDD:29008"
Site
                order (230, 240, 265, 267, 275)
                /site type="other"
                /note="ligand binding site"
                /db xref="CDD:29008"
                297..379
Region
                /region name="KR"
                /note="Kringle domain; Kringle domains are believed to
                play a role in binding mediators, such as peptides, other
                proteins, membranes, or phospholipids; cd00108"
                /db xref="CDD:29008"
Site
                309
                /site_type="other"
                /note="putative domain interaction site"
                /db_xref="CDD:29008"
Site
                order (324, 334, 359, 361, 369)
                /site type="other"
                /note="ligand binding site"
                /db xref="CDD:29008"
Region
                383..465
                /region name="KR"
                /note="Kringle domain; Kringle domains are believed to
                play a role in binding mediators, such as peptides, other
                proteins, membranes, or phospholipids; cd00108"
                /db xref="CDD:29008"
Site
                395
                /site type="other"
                /note="putative domain interaction site"
                /db xref="CDD:29008"
Site
                order (410, 420, 446, 448, 456)
                /site type="other"
                /note="ligand binding site"
                /db xref="CDD:29008"
```

//

```
490..714
     Region
                     /region name="Tryp SPc"
                     /note="Trypsin-like serine protease; Many of these are
                     synthesized as inactive precursor zymogens that are
                     cleaved during limited proteolysis to generate their
                     active forms; cd00190"
                     /db xref="CDD:29152"
     Site
                     490
                     /site type="cleavage"
                     /db xref="CDD:29152"
                     order (529, 573, 668)
     Site,
                     /site type="active"
                     /db xref="CDD:29152"
     Site
                     order (662,687,689)
                     /site_type="other"
                     /note="substrate binding sites"
                     /db xref="CDD:29152"
                     1..723
     CDS
                     /gene="HGF"
                     /standard name="hepapoietin A; scatter factor"
                     /coded by="M73240.1:66..2237"
                     /db xref="GDB:G00-127-524"
ORIGIN
        1 mwvtkllpal llghvllhll llpiaipyae ggrkrrntih efkksakttl ikidpalkik
       61 tkkvntadgc anrctrnkql pftckafvfd karkqclwfp fnsmssgvkk efghefdlye
      121 nkdyirncii gkgrsykgtv sitksgikcq pwssmipheh syrgkdlqen ycrnprgeeg
      181 gpwcftsnpe vryevcdipg csevecmtcn gesyrglmdh tesgkicgrw dhgtphrhkf
      241 lperypdkgf ddnycrnpdg qprpwcytld phtrweycai ktcadntmnd tdvplettec
      301 iggggegyrg tyntiwngib cgrwdsgyph ehdmtpenfk ckdlrenycr npdgsespwc
      361 fttdpnirvq ycsqipncdm shqqdcyrqn qknymqnlsq trsqltcsmw dknmedlhrh
      421 ifwepdaskl nenycrnpdd dahgpwcytg nplipwdycp isrcegdttp tivnldhpvi
      481 scaktkglrv vngiptrtni gwmyslryrn khicggslik eswyltargc fpsrdlkdye
      541 awlgihdvhq rqdekckqvl nvsqlvygpe gsdlvlmkla rpavlddfvs tidlpnygct
      601 ipektscsvy gwgytgliny dgllrvahly imgnekcsqh hrgkvtlnes eicagaekig
      661 sqpceqdygg plvceqhkmr mvlgvivpgr gcaipnrpgi fvrvayyakw ihkiiltykv
     721 pgs
```

Disclaimer | Write to the Help Desk NCBI | NLM | NIH

Aug 28 2007 16:53:42



```
JOURNAL.
            Cytokine Growth Factor Rev. 18 (3-4), 327-334 (2007)
   PUBMED
            17537668
  REMARK
            GeneRIF: FGF2 has a role in tumor angiogenesis associated with
            haematological malignancies [review]
            Review article
REFERENCE
            5 (residues 1 to 288)
  AUTHORS
            Arnaud, E., Touriol, C., Boutonnet, C., Gensac, M.C., Vagner, S.,
            Prats, H. and Prats, A.C.
  TITLE
            A new 34-kilodalton isoform of human fibroblast growth factor 2 is
            cap dependently synthesized by using a non-AUG start codon and
            behaves as a survival factor
  JOURNAL
            Mol. Cell. Biol. 19 (1), 505-514 (1999)
   PUBMED
            9858574
  REMARK
            GeneRIF: A new FGF2 isoform results from the use of a non-AUG (CUG)
            translation initiation codon.
REFERENCE
            6 (residues 1 to 288)
  AUTHORS
            Watson, R., Anthony, F., Pickett, M., Lambden, P., Masson, G.M. and
            Thomas, E.J.
  TITLE
            Reverse transcription with nested polymerase chain reaction shows
            expression of basic fibroblast growth factor transcripts in human
            granulosa and cumulus cells from in vitro fertilisation patients
  JOURNAL
            Biochem. Biophys. Res. Commun. 187 (3), 1227-1231 (1992)
   PUBMED
            1417798
REFERENCE
           7 (residues 1 to 288)
  AUTHORS
            Ago, H., Kitagawa, Y., Fujishima, A., Matsuura, Y. and Katsube, Y.
  TITLE
            Crystal structure of basic fibroblast growth factor at 1.6 A
            resolution
  JOURNAL
           J. Biochem. 110 (3), 360-363 (1991)
   PUBMED
            1769963
REFERENCE 8 (residues 1 to 288)
  AUTHORS
           Eriksson, A.E., Cousens, L.S., Weaver, L.H. and Matthews, B.W.
  TITLE
           Three-dimensional structure of human basic fibroblast growth factor
  JOURNAL
           Proc. Natl. Acad. Sci. U.S.A. 88 (8), 3441-3445 (1991)
   PUBMED
           1707542
REFERENCE
            9 (residues 1 to 288)
  AUTHORS
            Zhu, X., Komiya, H., Chirino, A., Faham, S., Fox, G.M., Arakawa, T.,
            Hsu.B.T. and Rees.D.C.
  TITLE
            Three-dimensional structures of acidic and basic fibroblast growth
            factors
  JOURNAL
            Science 251 (4989), 90-93 (1991)
   PUBMED
            1702556
REFERENCE
            10 (residues 1 to 288)
  AUTHORS
            Prats, H., Kaghad, M., Prats, A.C., Klagsbrun, M., Lelias, J.M.,
            Liauzun, P., Chalon, P., Tauber, J.P., Amalric, F., Smith, J.A. et al.
  TITLE
            High molecular mass forms of basic fibroblast growth factor are
            initiated by alternative CUG codons
  JOURNAL
            Proc. Natl. Acad. Sci. U.S.A. 86 (6), 1836-1840 (1989)
   PUBMED
            2538817
  REMARK
            GeneRIF: Alternate protein isoforms arise through the use of AUG
            and non-AUG (CUG) translation initiation codons.
COMMENT
            REVIEWED REFSEO: This record has been curated by NCBI staff. The
            reference sequence was derived from J04513.1, AC021205.6, M27968.1,
            BU501243.1, BP292299.1, CN315083.1 and AA256481.1.
            On Jul 24, 2007 this sequence version replaced gi:41352695.
            Summary: The protein encoded by this gene is a member of the
            fibroblast growth factor (FGF) family. FGF family members bind
```

heparin and possess broad mitogenic and angiogenic activities. This protein has been implicated in diverse biological processes, such as limb and nervous system development, wound healing, and tumor

growth. The mRNA for this gene contains multiple polyadenylation sites, and is alternatively translated from non-AUG (CUG) and AUG initiation codons, resulting in five different isoforms with distinct properties. The CUG-initiated isoforms are localized in the nucleus and are responsible for the intracrine effect, whereas, the AUG-initiated form is mostly cytosolic and is responsible for the paracrine and autocrine effects of this FGF.

Publication Note: This RefSeq record includes a subset of the publications that are available for this gene. Please see the Entrez Gene record to access additional publications.

FEATURES

Location/Qualifiers

gource 1.,288

> /organism="Homo sapiens" /db xref="taxon:9606" /chromosome="4" /map="4q26-q27"

Protein 1..288

/product="fibroblast growth factor 2"

/note="heparin-binding growth factor 2; prostatropin;

basic fibroblast growth factor bFGF" /calculated_mol_wt=30639

Region

/region name="24-kDa isoform; alternative non-AUG (CUG)

translation initiation site"

88 Region

/region name="22.5-kDa isoform; alternative non-AUG (CUG)

translation initiation site"

Region

Region

Site

CDS

/region name="22-kDa isoform: alternative non-AUG (CUG)

translation initiation site"

/region name="18-kDa isoform; alternative AUG translation

initiation site"

Region 163..285

/region name="FGF" /note="Acidic and basic fibroblast growth factor family;

FGFs are mitogens, which stimulate growth or

differentiation of cells of mesodermal or neuroectodermal

origin; cd00058"

/db xref="CDD:28940"

order (163, 166, 198, 200, 202, 230, 238, 241..246, 248, 280, 282, Site

284)

/site type="other"

/note="receptor interaction site"

/db xref="CDD:28940"

order (261..262,267,271,277)

/site type="other"

/note="heparin binding site (glycine box)"

/db xref="CDD:28940" 1..288

/gene="FGF2"

/coded by="NM 002006.4:69..935"

/exception="alternative start codon"

/GO_component="extracellular region [PMID 10490103];

extracellular space [PMID 2435575] "

/GO function="growth factor activity; heparin binding;

protein binding [PMID 11075807] "

/GO process="activation of MAPK activity [PMID 9712850]; angiogenesis; cell differentiation; cell proliferation;

```
cell-cell signaling; chemotaxis [PMID 10848592]; muscle
                     development; nervous system development [PMID 9576942];
                     organ morphogenesis [PMID 10903182]; positive regulation
                     of cell proliferation [PMID 2435575]; Ras protein signal
                     transduction [PMID 10848592]; regulation of progression
                     through cell cycle; signal transduction [PMID 9712850] "
                     /note="non-AUG (CUG) translation initiation codon; 34-kDa
                     isoform"
                     /db xref="CCDS:CCDS34059.1"
                     /db xref="GeneID:2247"
                     /db xref="HGNC:3676"
                     /db xref="HPRD:00622"
                     /db xref="MIM:134920"
ORIGIN
        1 mvgvgggdve dvtprpggcq isgrgargcn gipgaaawea alprrrprrh psvnprsraa
       61 gsprtrgrrt eerpsgsrlg drgrgralpg grlggrgrgr apervggrgr grgtaapraa
      121 paargsrpgp agtmaagsit tlpalpedgg sgafppghfk dpkrlyckng gfflrihpdg
      181 rvdgvreksd phiklqlqae ergvvsikgv canrylamke dgrllaskcv tdecffferl
      241 esnnyntyrs rkytswyval krtqqyklqs ktqpqqkail flpmsaks
//
```

Disclaimer | Write to the Help Desk NCBI | NLM | NIH

Aug 28 2007 16:53:42

Blast Result Page 1 of 1

Structure



DI ACT 2 SECUENCES DESIII TS VEDSION DI ACTD 2 2 17 IAug 26 2007

BLAST 2 SEQUENCES RESULTS VERSION BLASTP 2.2.17 [Aug-26-2007]									
~ *	ect: 10.0000 wordsiz ion X for protein, n f	ap extension: 1 ae: 3 <u>Filter</u> View of View of Masking		4					
Sequence 1: unnamed Length = 1207	protein product E	GF							
Sequence 2: unnamed protein product HGF Length = 723									
No significant similar	ity was found								
CPII time: 0.03	user secs.	0.02 sys. secs	0.05 total secs						

Blast Result Page 1 of 1

Structure

0.05 total secs.



PubMed Entrez BLAST OMIM Taxonomy

BLAST 2 SEQUENCES RESULTS VERSION BLASTP 2.2.17 [Aug-26-2007]

Matrix BLOSUM62 gap open: 11 gap extension: 1								
x_dropoff: 0 expect: 10.0000 wordsize: 3 Filter View option Standard								
Masking character option X for protein, n for nucleotide Masking color option Black								
Show CDS translation Align								

Sequence 1: unnamed protein product EGF Length = 1207

Sequence 2: unnamed protein product FGF-Z Length = 288

No significant similarity was found

CPU time: 0.04 user secs. 0.01 sys. secs

Blast Result Page 1 of 1



PubMed Entrez BLAST OMIM Taxonomy Structure

BLAST 2 SEQUENCES RESULTS VERSION BLASTP 2.2.17 [Aug-26-2007]

Matrix BLOSUM	62 gap o	oen: 11 gap	extension: 1				
x_dropoff: 0	expect: 10.0	000 wordsize:	3 Filter	☐ View option §	Standard		
Masking charact	er option X f	or protein, n for	nucleotide ·	Masking color	option B	lack	
Show CDS tra	anslation [dign					
		_					
Sequence 1: unn Length = 723	amed proteir	product HG	aF				
Sequence 2: unn Length = 288	amed proteir	product F	4F-Z				
No significant si	milarity wa	s found					
CPU time:	0.03 user	secs.	0.02 svs. s	secs	0.05 to	tal secs.	